Nucleotide sequence of SSL2 genomic gene (wild type)	
A T G A G T A G T T T G G T G G A G G C T T C G C A T A C G A T C T G A T A G G A A A C C A G T T T A T A A C C T A G A T G A T C T G A T G A T G	9 2
A C G A C T T C G T T C C T A A A A A A G A T C G A A C C T T T G A G C C A A G T C G A G G C T A T T G T G A G G E T T T G A G C C A G G C T A T G T G A G A C T G A T G C G G T T T G A C C C A G A G C T A T G C A G A C T G A G G C T T T G A G C A G G C T A T G T G A G A C T G A T G C G G T T T G A C T G A G A C T G A T G C A G A C T T G A G A C T G A G A C T G A G A C T G A G A C T G A G A C T G A G A C T T G A G A C T T G A G A C T T G A G A C T G A G A C T G A G A C T G A G A C T G A G A C T G A G A C T G A G A C T G A G A C T G A G A C T G A G A C T G A G A C T G A A G A C T G A G A G A C T G A G A G A C T G A G A C T G A G A C T G A G A C T G A G A G A C T G A G A	152
ctctcgagcttattgttcagcttttactgttttatgtgtgttctatttaatctattttagtctttttttt	2 2 8
tttgtagAAAGAAAATGCATGTCAGGCTTGTGGGGAAAGTACTAATCTTGTAAGCTGCAATACATGCACTTATGCG	304
TTCCATGCTAAATGCTTAGTTCCACCTCTTAAAGATGCTTCCGTGGAAATTGGAGATGCCCTGAATGTgtaagat	3 8 0
tttagttacggtccacaattatgttttgggatgctacaggttccatttttttt	456
t t t g c a g G T T A G T C C T C T T A A C G A G A T A G A T A T T G G A T T G T G	532
G G T T C C T C C G A T G C G G A A G C C A A T T T T T G T G A A C A G T A T C T C G T G A A G T G G A A G G G A T T A T C A T A C C T T C	809
ACTGCTCTTGgtagttactgcgtgtcttttttgctgtctggacacgctaattatcaatgtttctttc	684
tataatatgtgatttatttccttttactaatcatagGGTGCCTGAGAAGGAGTTCCAGAAGGCTTATAAGTCAAAT	260
CAT C G T T T A A A A A C C A G A G T G A A C A A T T T T C A C C G T C A A T G G A G T C A T T C A A T A A C A G C G A A G A T G A T T T T G T T G	836
CCATACGTCCTGAGTGGACCACTGTTGATCGGATTCTTGCCTGCAGgtctagagaatggaattaatttattt	912
atctatctgccaactttttttttaatatccttgttttcagcataatccattctttaataaacacgtatctttgata	8 8 6
gagtgetgettaacetaaatttactgttatcacgattttgggtetetgaaacatgataaagtgactgaett	1064
ttttcttcttcttttaagttaccattttcttagttgtttcgtaaatcaggaattgtgtgcattggcattggttcttt	1140
tatgatatagAGAAGAAGATGGGGAGCTGGAATATCTTGTCAAATATAAAGAGCTATCCTATGATGAAGTGTTATTG	1216
GGAGTCAGAATCAGACATCTCAACCTTCCAGAATGAAATTCAAAGGTTCAAGGATGTAAATTCTAGAACTCGCAGA	1292

<u>ئ</u>

A GTA A A GATGTT GACCATA A A A A A A A TCCCA GA GACTTT CA A CAGTTT GATCATA CTCCT GAATT CCT CAAA G g t a	1368
tttggatcaccttaaatcatatactataaatgtttcttatatttggtacttatagatgttatgatttattgtttg	1444
	1520
TGTAATCC	1596
catccattgtaaggggcctttgtttctattcctgtaatgttgtgagattttcctgttacagGCAAGACAATTCA	1672
A A G C A T T G C C C T T T T T A G C T T T T T T G A G A G A G A A C C T C A T T C C G C A T T T G G T A A T T G C T C C T C T A T C G A C T C T G	1748
CGTAACTGGGAGAGAGATTTGCCACATGGGCCCCACAGATGAACGTGgtatgtatgcagttatacacgcaatgat	1824
ctgtgccatttgtatgtttttgtttgtttgttaatggaatggtcttcgtggtcatttgacgggtagGTTATGTATTTT	1900
T G G C A C T G C G C A G C T C G A G A T T A T C A G A G A C A T G A G T T T T A C T T A T C G A A G A T C A A A A A A A A A A A A A A A A A	1976
A A G A A A T C T G G A C A A A T A A G C A A A G C A A A A A A A A A A G A A G T T T G A T G T C C T C C T C A C A T C G T A T G A G A	202
TGATCAACCTAGATTCAGCAGTTCTAAAACCAATTAAGTGGGAGTGCATGgtaactcttattctctaatgagactt	2128
tactttctcttagtcgtctctttctctcttacatgttgcctagtaacaattgtttggggcagATTGTTGATGAA	2204
GGTCATCGACTGAAAAATAAGGATTCAAAGCTGTTCTCTTCATTGACACAGTATTCAAGTAACCACCGTATTCTTC	2280
TGACAGGAACACCACTTCAGgttcgtcatttgagtttgatttctgaagtttatactttcaatagttgtatctgagc	2356
atagtagctacgatttgcaatgagaattgttatatattatcttgcactaatgtcttacctgattagttgcaatatg	2432
ttactgatgattatgtggtgcctttacagAACAACTTGGATGAACTTTTCATGCTCATGCATTTTTTGATGCGGG	2508
GAA G g t a t c a c a a g a a t a g a t a a a t a a g t t c g c a t a c t t a a c a g a a t t t t a t g t a g c t a a c a t g t t a t t t g	2584
attgcacaatacttgcagTTTGGAAGTTTGGAGGAGTTCCAGGAGGTGCAGAAGATATTAATCAAGAGGAGCAGA	2660

F | G. 2

T C T C A A G G T T G C A C A A A A T G T T G G C T C C A C A T T T G C T C A G A A g t a t t a a a c a a a a c t a t t t g t t c a t c t t t t t t a a	2736
tttatatatattcaaaagtttggttggagggaatctttcatagtaatatttatgatcttatgatcttaaccatgctgtctc	2812
tattttgattgctcttccagGGGTAAAAA	2888
GTTGATCTGAGCAGTCTGC	2964
GGAGGTGCTCAAgtaagttctttttaatttttgtttacactttttggatcattaaaacctcataggtggggtagaaa	3040
a a t	3116
ATTACGAAAAGTATGCTGCCATC	3192
AAqta	3268
a a c a t t t t c t t a t a t g t a t c a a a g A C	3344
AAACTGAAAGAGCAAGGACACAG	3420
TGTACCCATAAGgtatttgaacttcttatatgtacagtctgtttcagtagaattttcattcttgttgttttgtaga	3496
a c t g t	3572
aaatt	3648
acaqtacaaatcacaagt	3724
gtatacagagtteteat	3800
gaacttgt	3876
cactc	3952
ttctt	4028

7 ---

qattqctcctttcttatgaaagcttttgcagtcaattgcatggggcgtatttcatttgtctctatcttctgtct	5472
gcagATTGCTTGATCGTGATCTCGTGGAG	5 5 4 8
ر و ر	5624
caatgetecattggagtetatgettaattgtattettatattecatgatatteagaetggeggetgggttggetategaaat	5700
ctgctgtg	5776
ىد ب	5852
ļ	5928
tttgaaccctgatc	6004
tgcttg	0809
gttctgaagcaa	6156
atcttcc	6232
CGCTGCTGAAAGCAAA	6308
TTGAGCTGCACCAGGCTGAGGAGCTTAATGCTCTTGGAAAAAGGAAGAAAGGCAAGTCGCAAGCAGG tttggtctct	6384
t gat c c c c c t t a t c c a a t t g t g g c a t c a t a t t g a t a a c t g a t t t t t c a t t t a t g t t c t t t c t g a t t c t g t c c	6460
t gett catatat ttatt cat get get cta a cett te cett tt ga a tt cett a g g ta g cta a a tt caga a a g ta a ta a	6536
t tag t t gact g tatc ctt c taaat t gagaaag tataat t tag t t gact g tatc cag tataaaac taaac g c c c t t g	6612
Ü	8 8 9 9
ttaactatagtettetgttteetetgeaagaaataegttttgttteaetetetaaettgatatageteegeteaattaet	6764

5

tacattggtttggtctgccatcatcgtttcatgtcttcaatuataggctgctgctgctgctgtccaattagg
GGTATCCAT
G A G T C A A C A G A T G G T G A A G C A G C A G G A C A A G G A G T T C A G A C G G G T C G A C G T A C A G A A A A
t taccacg t t t c g g a t t t a a t t t a a t t t g t a a t g g a g c t g a a a t g a c t g a t a t t a g a a g t g t g c
tttt
GTTTCAACCAGAGTCAAA
cg c c a g a a g c t a t t c t g t c t a a a t t t t a c a g t t t c a t c c c c c g a t g c a t c t a a a t t a t c a t c a g t c
atatt
TTCCTCGCTTAAAGCAGAAGACCTTTGAAGAAATAAATGAgtacgggctcaaccctttaatgctct
t t c t t t a c a a a a a a c g c a t c a t t a t a a a a a g g c t t t c t g g t t t a t t t t t a a t t t t t t
a g A T A T G G A A T A C T C T T C T T G A G C A C A T T G C T G A A G A A A T A G A C G A G A A T T C T C C A A C C T T T T C A
a a t t g a t a t t t t c a c t g t t t g c t t t t t c c c t a a a t g a g a t c a t t g c t t c t c t g t t a a c c g g t t
atataatggtcgttgtctatagATGGTGTCCCAAGGAAGGACTTAGAATAGAAGGTGTTCTAGTC
TTCTGATACTAGTTCAGGAGAAGgtgagtctattgactttaattcttcattaagttctctttta
ttttttggtatatgttacttctagtctatagtttagctctgttcgcataagtttttaatacagtaat
a a g a t t
T C A T C C A G G G A A A C C T G T T T T T C C C C T C T C G C A T T C T T G A A G A T T C C C C G G A C T G A G A G T G G A A A
G A G G A A C A T G A C A A G A T A A T G A T A C G T G C T G T T T T A A B t a t g a a c c c t g c a c c a c t

aatggttttatttctcatcattctccattacttgctcacattttctttc	8 2 0 8
T C C C T C A C A T A A G T T T G T C T G C T G A A G A G C T G G T T T G C A G G G G G G G G T A G T G G T A G T G G G G	9 e
C C G G G G G G C A C A G A A C C A G A A T C C T G G A A G C G T T A T T A C T G G A A C A A T A A T G C T T C T G C T G A T G G G G C T C A A G	8436
T A A A C T C G A T G T T C T A T T A T C G G G A C A T G C A G A C G A C T T G T T G A G T T G T G A A A A A G C G A G T T C T G C T T T T G G A	8512
GAAGGCGATGAATTATGAATACGCAGAGGAATATTATgtatgttgtaccatctgcagtgttggtacttactcacat	8 5 8 8
gttttgcgctgaattgtttaactttgattgaatctctggttgcagGGACTTGGTGGCCTCATCATCTATCCTACTG	8664
A A G A A C C A G A A G C T G A A G A T C G C T G A C A C A G T G G G A G T G A G C T T T A T T G A G G T T G A T G A T G A A A T G C T T G A	8740
T G G A C T T C C T A A G A C T G A T C C T A T C A g t a a g t t c c a t c a c a a g t t t t a t t t a a c g a g t t g t t c t a a t g t	8816
gagetetetgaatetegetgeagCTTCAGAAGAAATTATGGGGGCTGCTGTTGACAACAACCAAGCGGGGTGGAA	8 8 9 2
ATAGCTCAACATTATAACCAGgtaagctatgctttttcctttggtggtaggctaatgtctagaactagtatatca	8968
cactaatatctctccggttattcagATGTGCAAACTTCTTGATGAGAACGCTCGGGAATCAGTCCAAGCATATGTA	9044
A A C A A C C A A C C A C C A G G T G A A T G A G C T T C C G T G C A C T C A A T C T A T C A A T G G T A A C A T T A A C A C A A	9120
T C C T T T C G A T T A C A T C T G A T C C A A G T C A C A T G A A G A C G A C C A A G C C A G A C C T A A A C A A T G T T G A G A T G A A	9196
GGACACGGCCGAAGAAACAAACCGTTAAGAGGTGGCGTCGTCGATCTGAATGTGGGGGGGG	9272
GCTGAAGCTAGTGGAAGTGTTGATGTAAAATGGAAGAAGCCAAAGAAGAAGAAGAGGGAGAGCCAAAGAACATGGTCGTTG	9348
ATTGA	9353

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Nucleotide sequence of mutant IAA14 gene	
ATGAACCTTAAGGAGAGGGCTTTGTCTTGGCCTCCCGGAGGCACTGAAACCGTTGAAAGTCCGGCAAGTCGG	9 /
GTGTTGGGAACAAGAGAGGCTTCTCCGAGACCGTTGATCTCAAACTTAATCTTCAATCTAACAAACA	152
GGATCTCAACACTAATGGAGCTCCCAAGGAGAGACCTTCCTT	228
ctatttacacaattccttaagaagaagaccttccttaaaagggaagactttttttt	304
actoatagttgatataaaagttettaaaataeatatatatatgaaagatgtaaggatgeataeagtaataaegttattg	3 8 0
a a t g t g t g t g t t g t t a t a t t c t a t g c a g A G C A C A A G T G G T G G G T T G G C C A T C G G T G A G G A A C T A C C G G A A A A	456
A T G T T A T G G C T A A T C A G A G G G G G G G A G G A G G A G G C A A T G A G T G G T G G G G A A C C G T C G C C T T T G T G A A	532
G G T T T C C A T G G A T G G A G C T C C T T A T C T T C G G A A G G T T G A C C T C A G A T C T C T C T C T C T C T G A T	608
GCCTTGGCCAAAATGTTCAGCTCCTTTACCATGGgtatgcattttcagacatataagtcgaattatcattatt	684
tttgtgtttacttacaatttttttttttaacgatacagttttttccatatacgactaattaat	092
gggattttgattaattaag GGAGTTAT GGAGCACAAGGGATGATAGATTTCATGAACGAGAGTAAAGTGATGGATC	836
T G T T G A A C A G T T C T G A G T T C C A A G C T A C G A G G A C A A G A T G G T G G A T G C T C G T T G G T G A T G T C C C C T G	912
GCCgtgagtttcctcattcttcttgctttcattattatgaccaaaattattctctaaacaaaaaaaa	9 8 8
ctaaagcattattattgatattacttatcaaaaaatacacaaaatgataatcaattatcatgtgttataaacacg	1064
cacagecatettttggttggeatgggaeagaaeteagagaeagageagagaegatgtttatatata	1140
atatgttacctcatttgtagctggcacatattctttcactttcaatagatttctaaatttagtcaccaacccaaat	1216
c c c g a t t t c a g G A T G T T G T C G A G T C A A C G T T T G C G C A T A A T G A A G G A T C C G A A G C A A T T G G A C T T G g t a	1292
agttttcttttctgttcgtttctataagtggctcttttctgttttccaataatgctcgtgtttttttt	1368
CAAGAGCAATGGAGAAGTTCAAGAACAGATCATGA	1403